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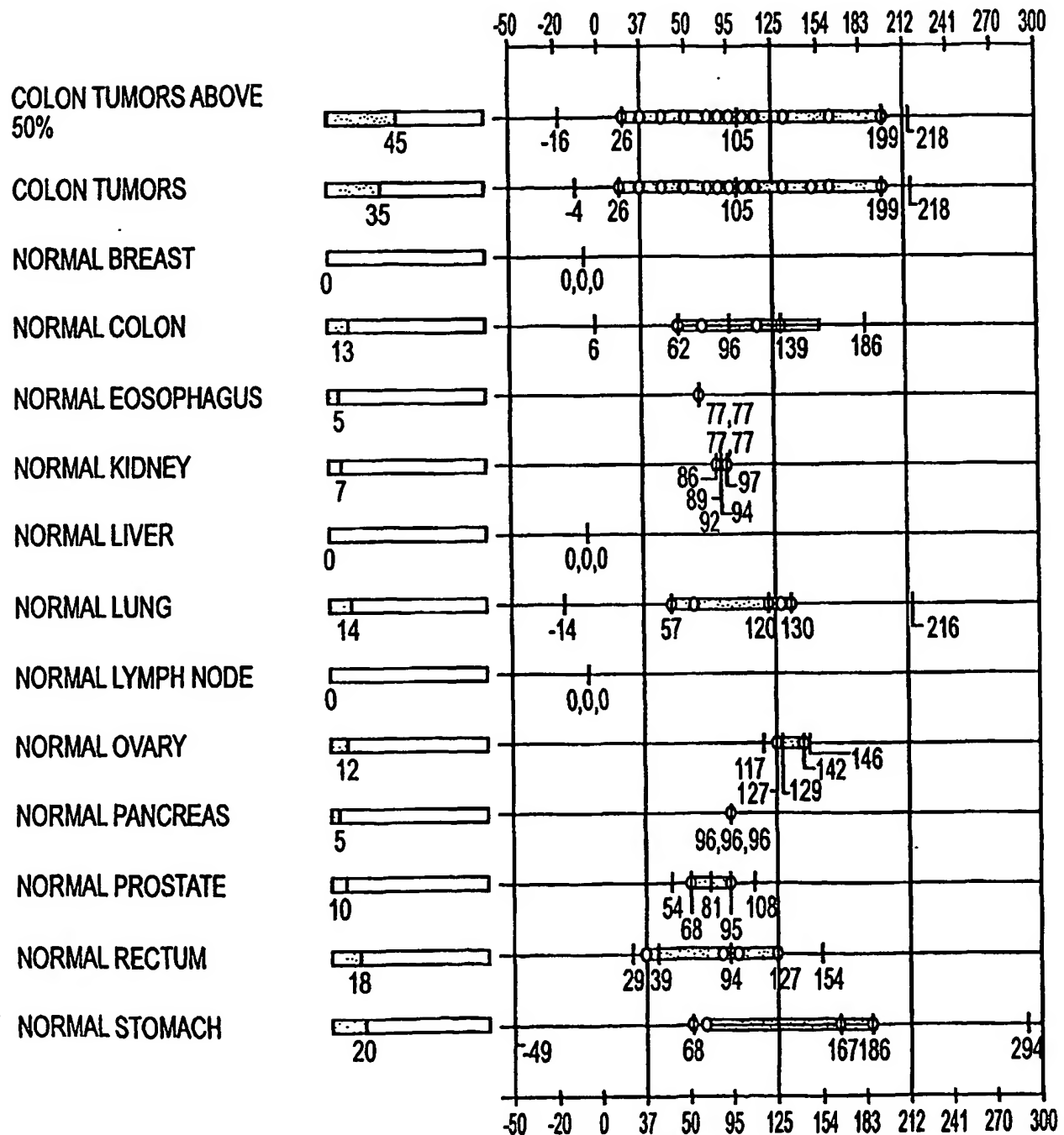
NAME	DESIGNATION	SIZE	A TUMOR	A TUMOR	AN:N	ANORMAL	BT:N	B TUMOR	POOL	POOLED NORMAL	DT:N	D TUMOR
CIC01	BS210MS130	184.75	3	1213.04	3	256.05	3	2685.32	2	298.09	3	1812.48
CIC02	BS220MS230	191.20	3	994.12	3	108.37	3	228.49	3	60.10	3	1271.98
CIC03	BS430MS430	222.32	3	1587.80	3	113.79	3	1226.02	3	117.06	3	163.29
BS421MS433												
		258.00	3	213.22	3	62.02	3	175.93	2	36.16	3	158.55
FT2:N	F TUMOR	FN2:N	F NORMAL	GT:N	G TUMOR	GN:N	G NORMAL	HT:N	H TUMOR	HN:N	H NORMAL	IT:N
3	658.97	3	229.01	3	2750.16	3	307.80	3	858.10	3	271.62	3
3	370.06	3	41.80	3	590.73	3	56.69	3	295.42	3	90.93	3
3	2747.40	3	118.80	3	1559.78	3	112.18	3	302.67	3	66.51	3
3	148.52	3	45.92	3	306.88	3	115.65	3	168.77	3	75.19	3
IN:N												
	I NORMAL	MT:N	M TUMOR	MN:N	M NORMAL	QT:N	Q TUMOR	QN:N	Q NORMAL	RT:N	R TUMOR	RN:N
3	225.34	3	3354.80	3	267.40	3	1494.28	3	261.62	3	2518.73	3
3	57.46	3	90.37	3	47.53	3	132.73	2	66.31	3	376.15	3
3	90.56	2	280.45	3	245.67	3	2593.52	3	158.56	3	1061.67	3
3	29.94	3	95.06	2	34.97	3	237.60	3	24.70	3	293.87	3
8TN												
	8 TUMOR	TUMOR AVE	NORMAL AVE	AVE FOLD	# OF HITS							
3	256.60	1827.61	296.41	6.17	20							
1	130.50	513.22	64.94	7.90	35							
3	160.96	1790.40	121.49	14.74	19							
3	248.80	203.42	53.11	3.83	22							

FIG. 1

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AFFY NAME : 52304\_AT

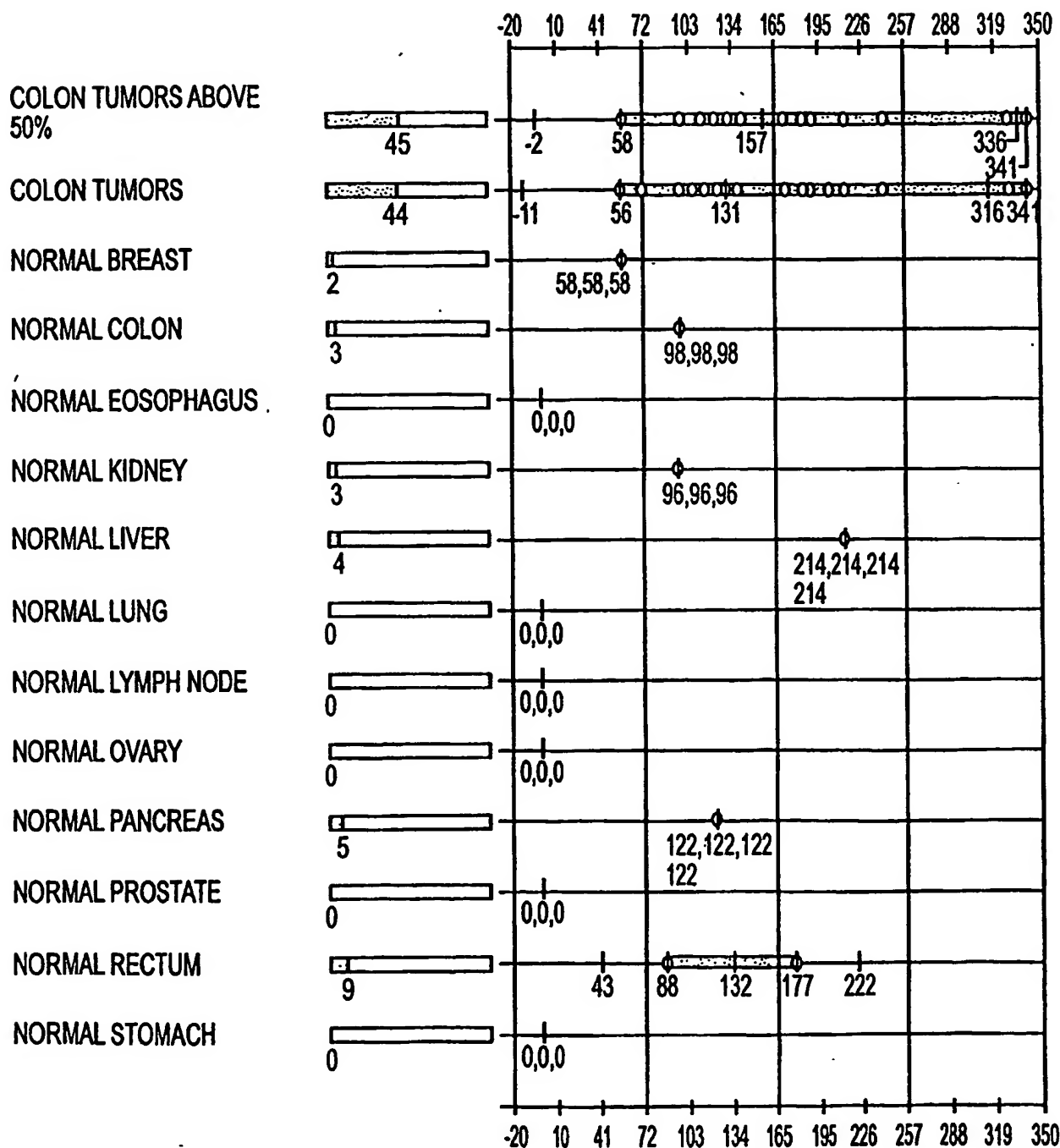
CLUSTER NAME : ESTs

**FIG. 2**

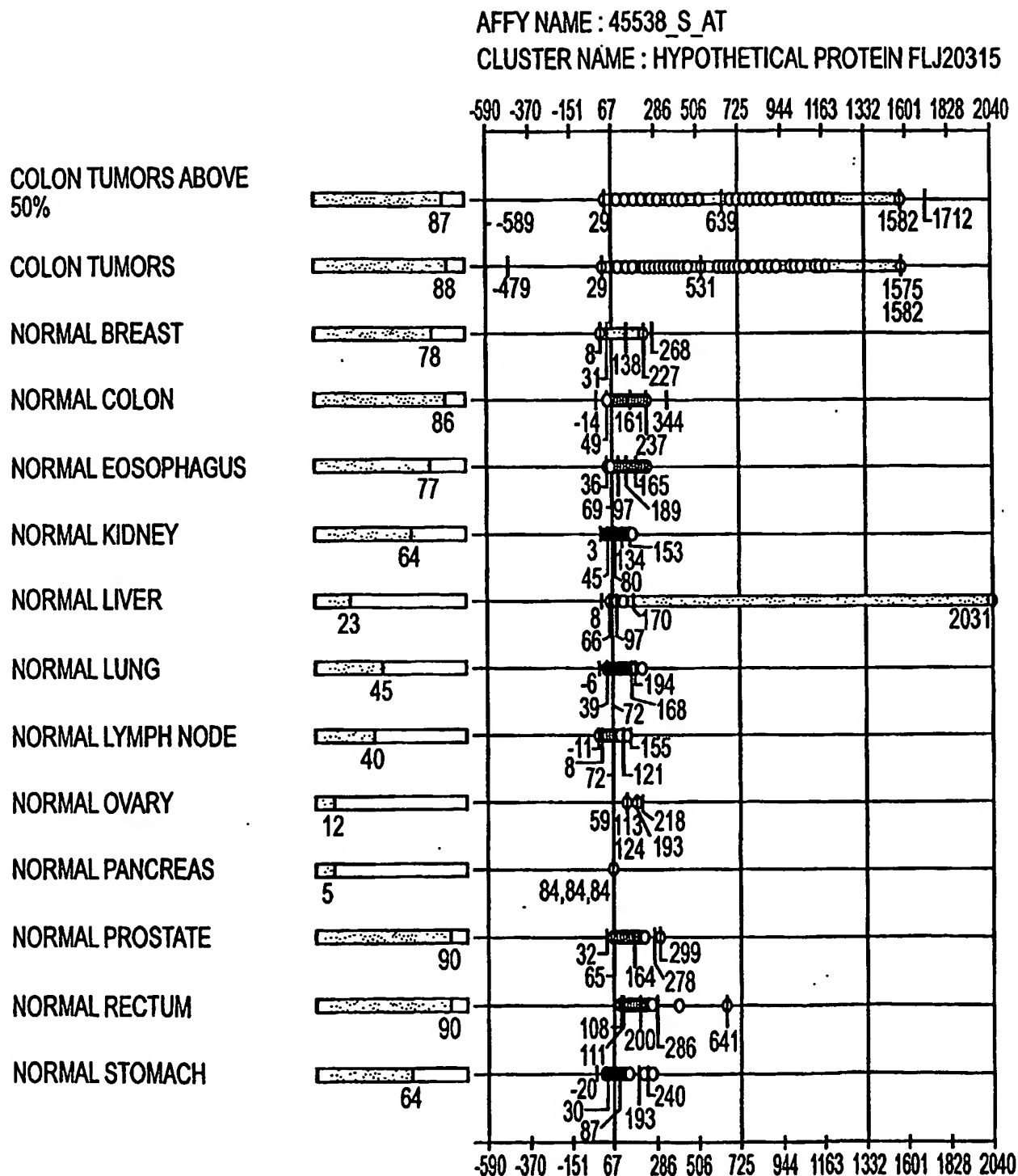
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AFFY NAME : 57471\_AT

CLUSTER NAME : ESTs

**FIG. 3**

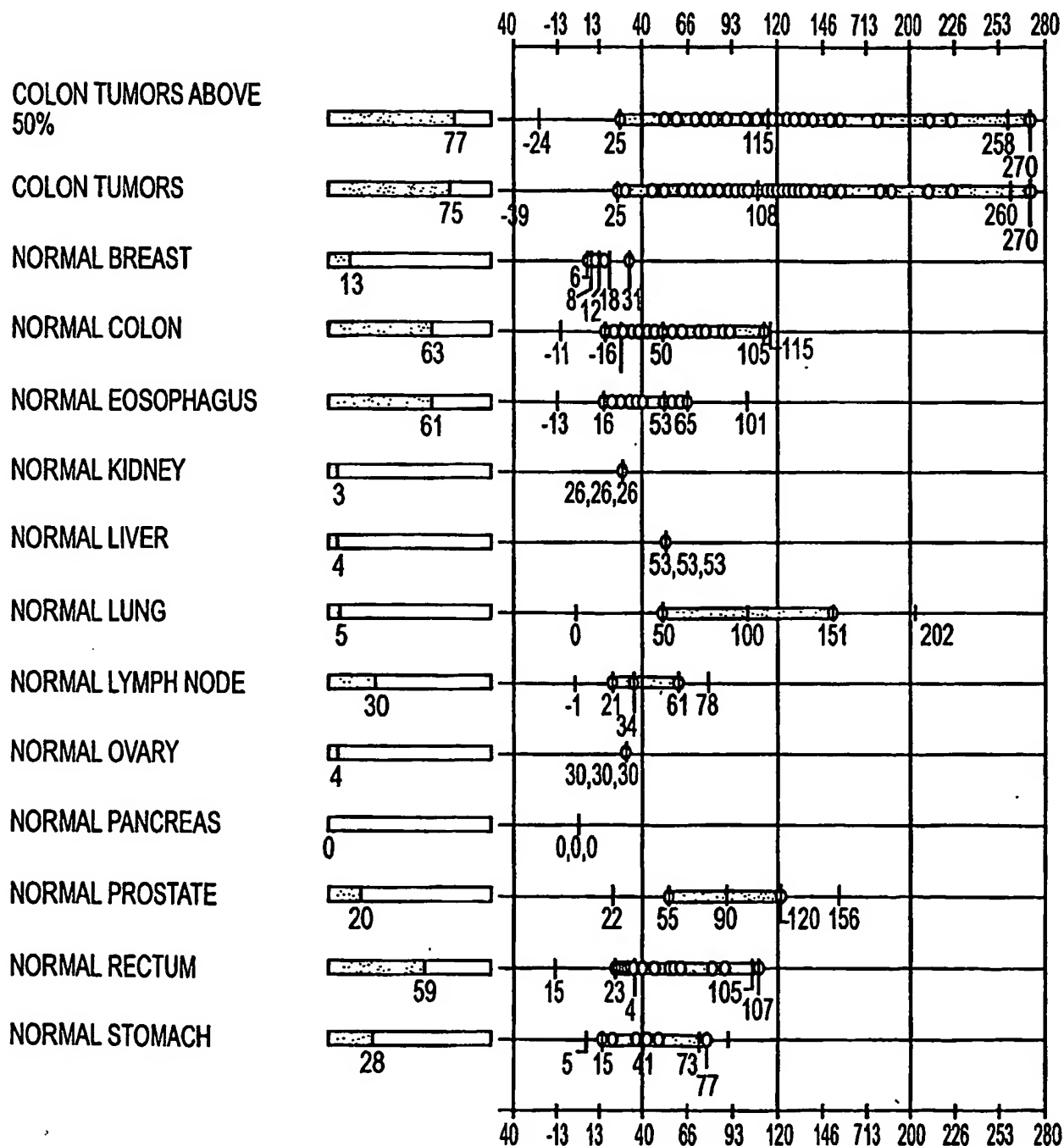
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**FIG. 4**

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AFFY NAME : 56634\_AT

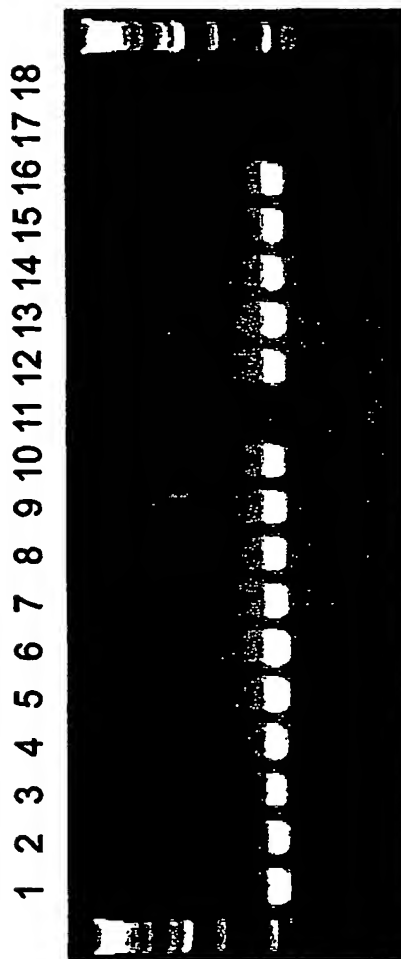
CLUSTER NAME : HYPOTHETICAL PROTEIN FLJ20354

**FIG. 5**

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**FIG. 6A**



**FIG. 6B**

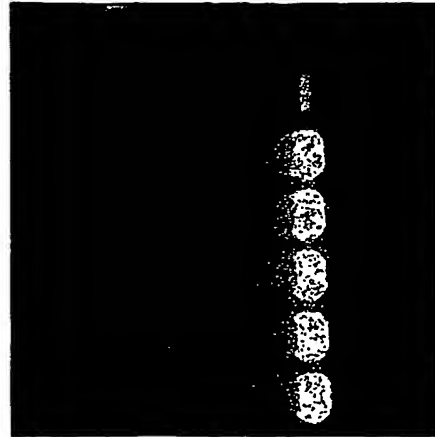
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1  
2  
3  
4  
5  
6  
7



**FIG. 7A**

1  
2  
3  
4  
5  
6  
7

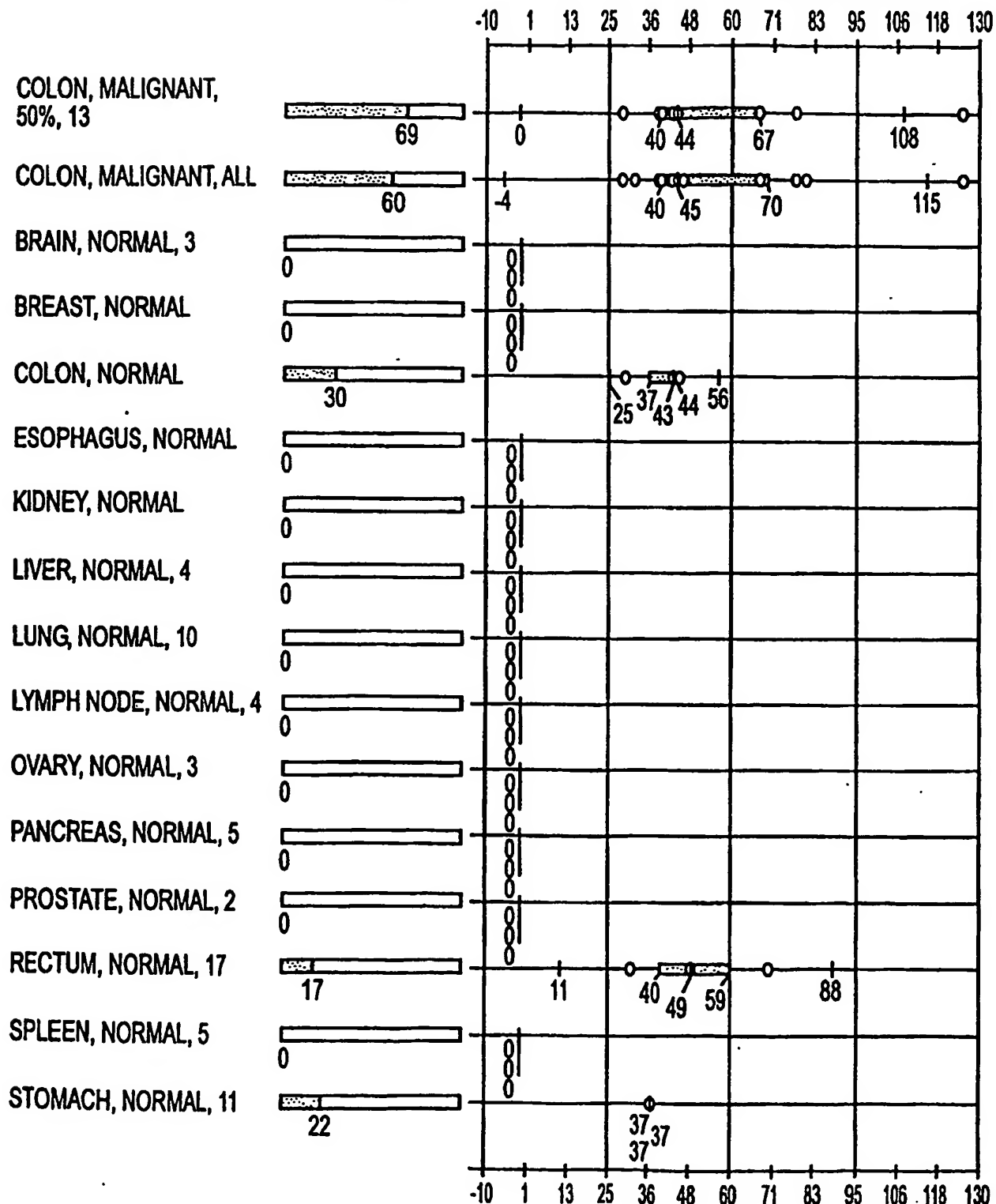


**FIG. 7B**

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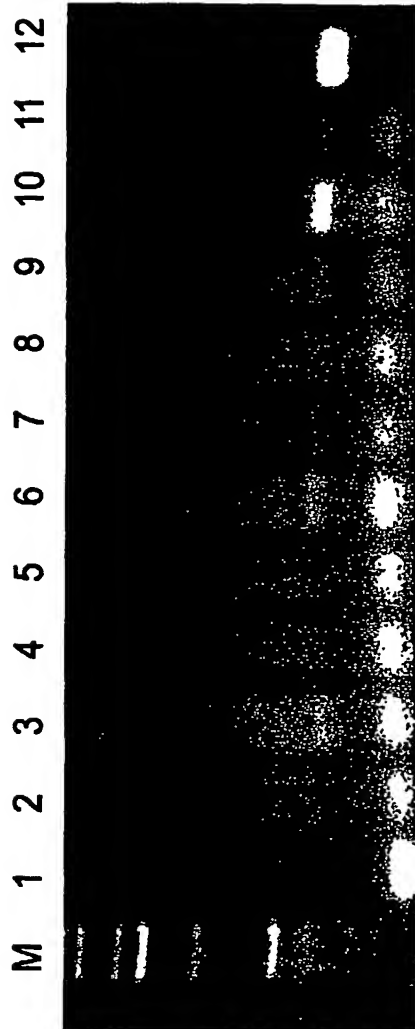
AFFY NAME : 222206\_S\_AT

CLUSTER NAME : HYPOTHETICAL PROTEIN FROM EUROIMAGE 2021883

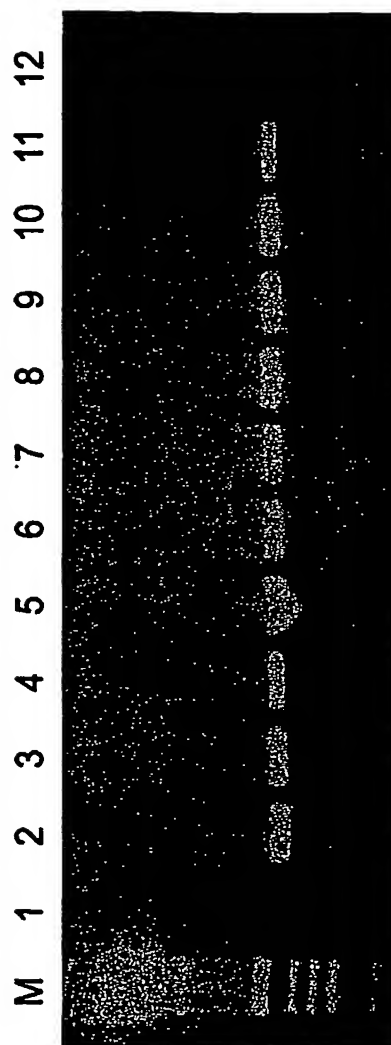
**FIG 8**



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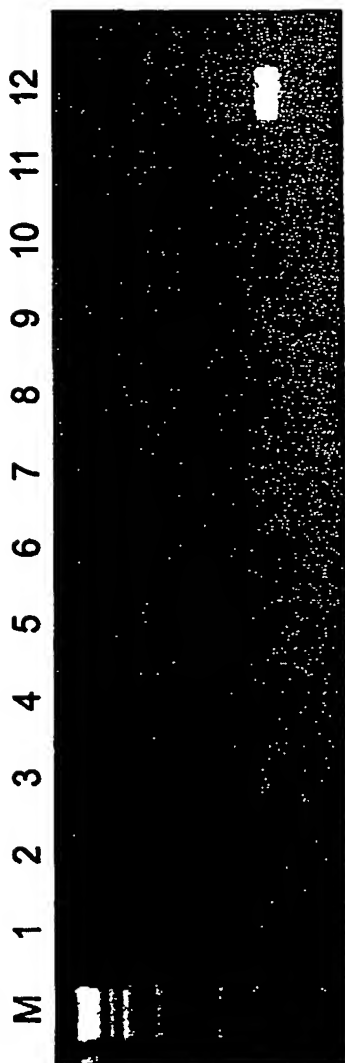


**FIG. 9A**

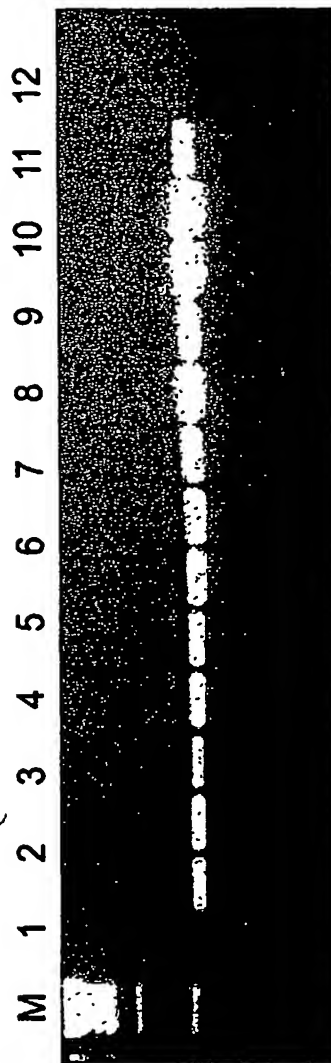


**FIG. 9B**

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**FIG. 10A**



**FIG. 10B**

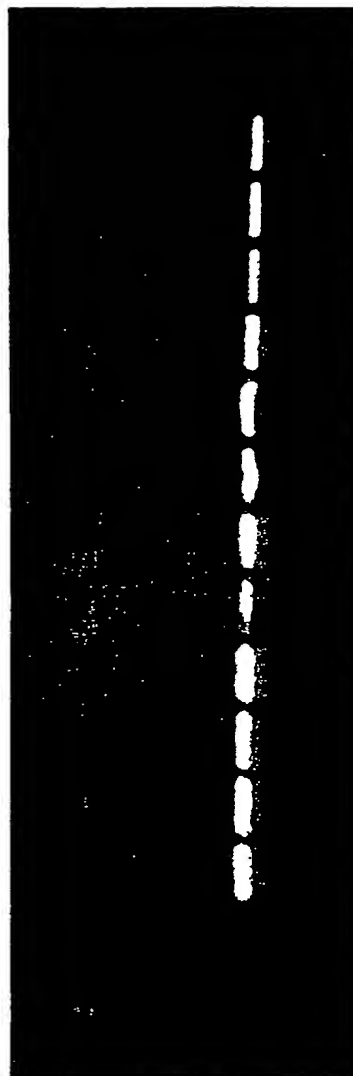
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M 1 2 3 4 5 6 7 8 9 10 11 12 13 14



**FIG. 11A**

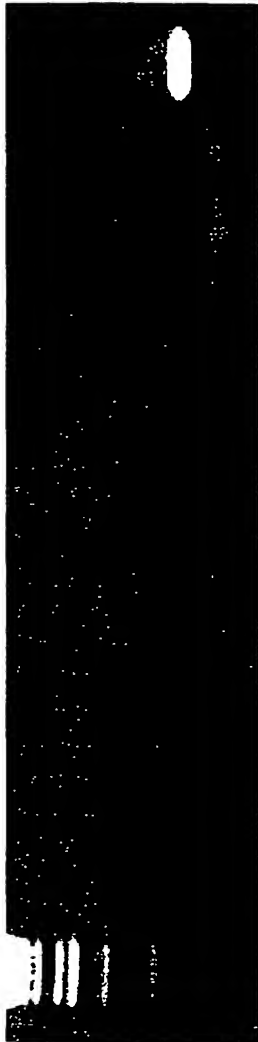
M 1 2 3 4 5 6 7 8 9 10 11 12 13 14



**FIG. 11B**

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M 1 2 3 4 5 6 7 8 9 10 11



**FIG. 12A**

M 1 2 3 4 5 6 7 8 9 10 11



**FIG. 12B**

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AFFY NAME : 238567\_AT

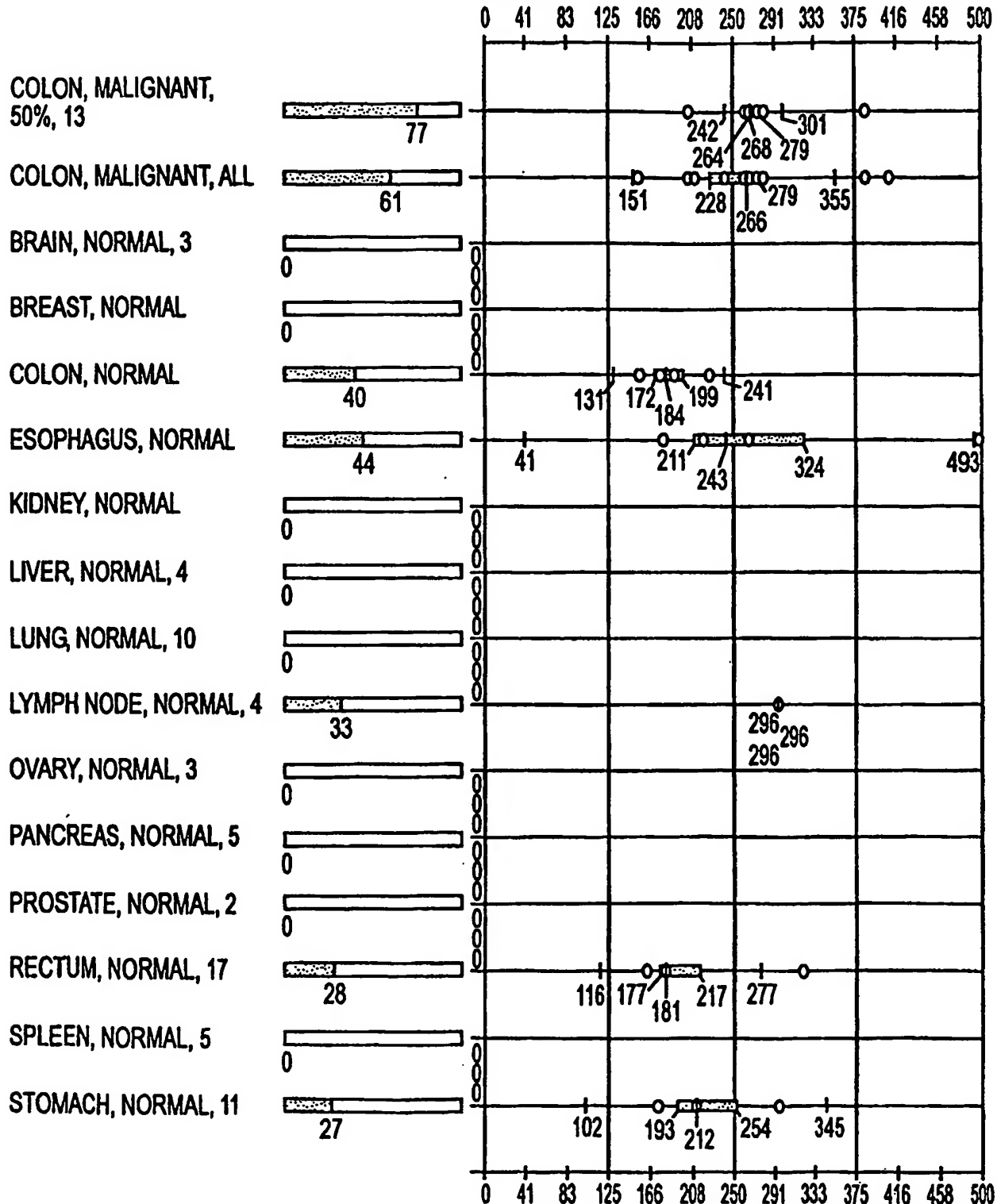
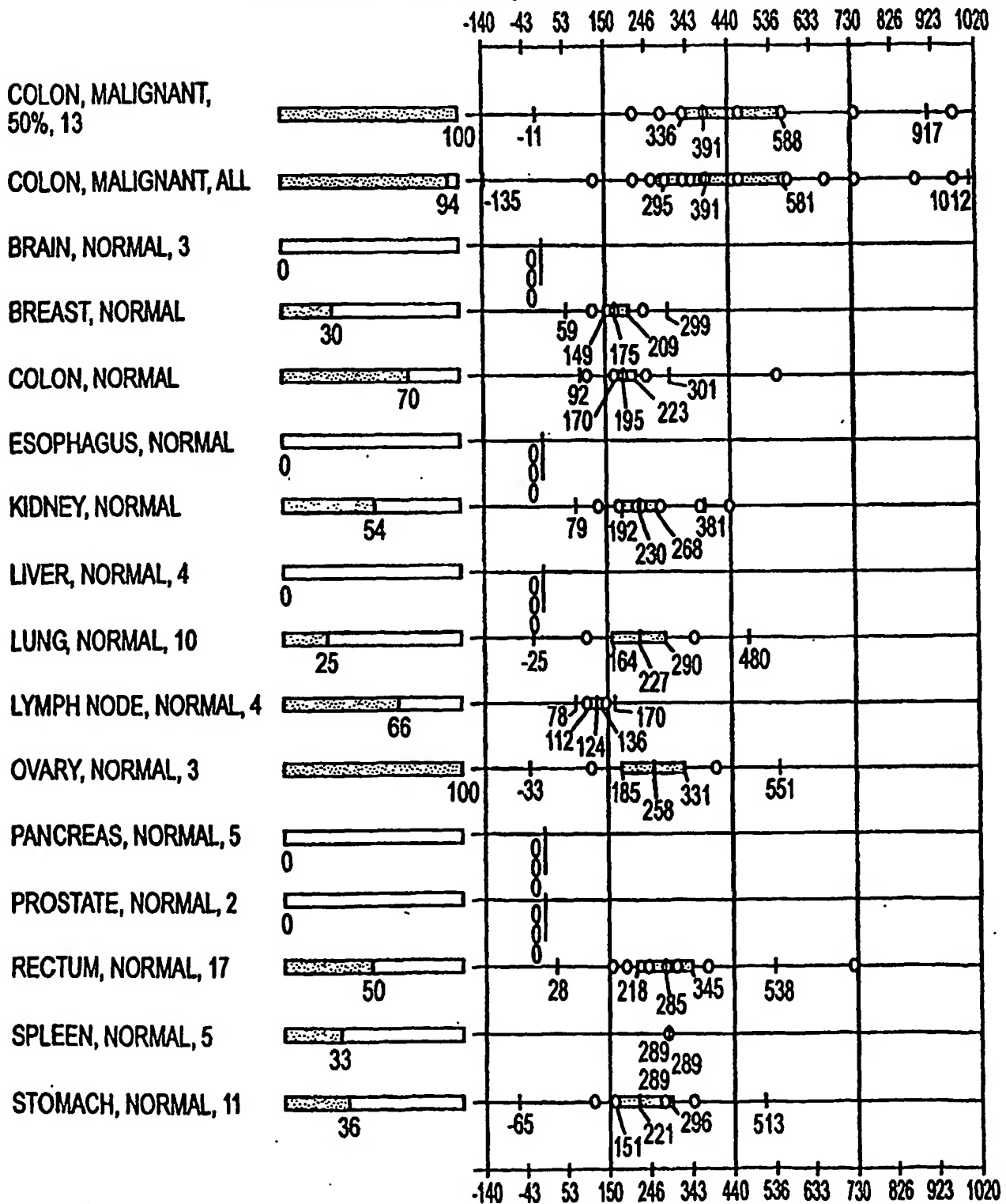
CLUSTER NAME : ESTs, WEAKLY SIMILAR TO YK33\_YEAST HYPOTHETICAL  
46.5 KD PROTEIN IN MRS4-DYN1 INTERGENIC F

FIG 13

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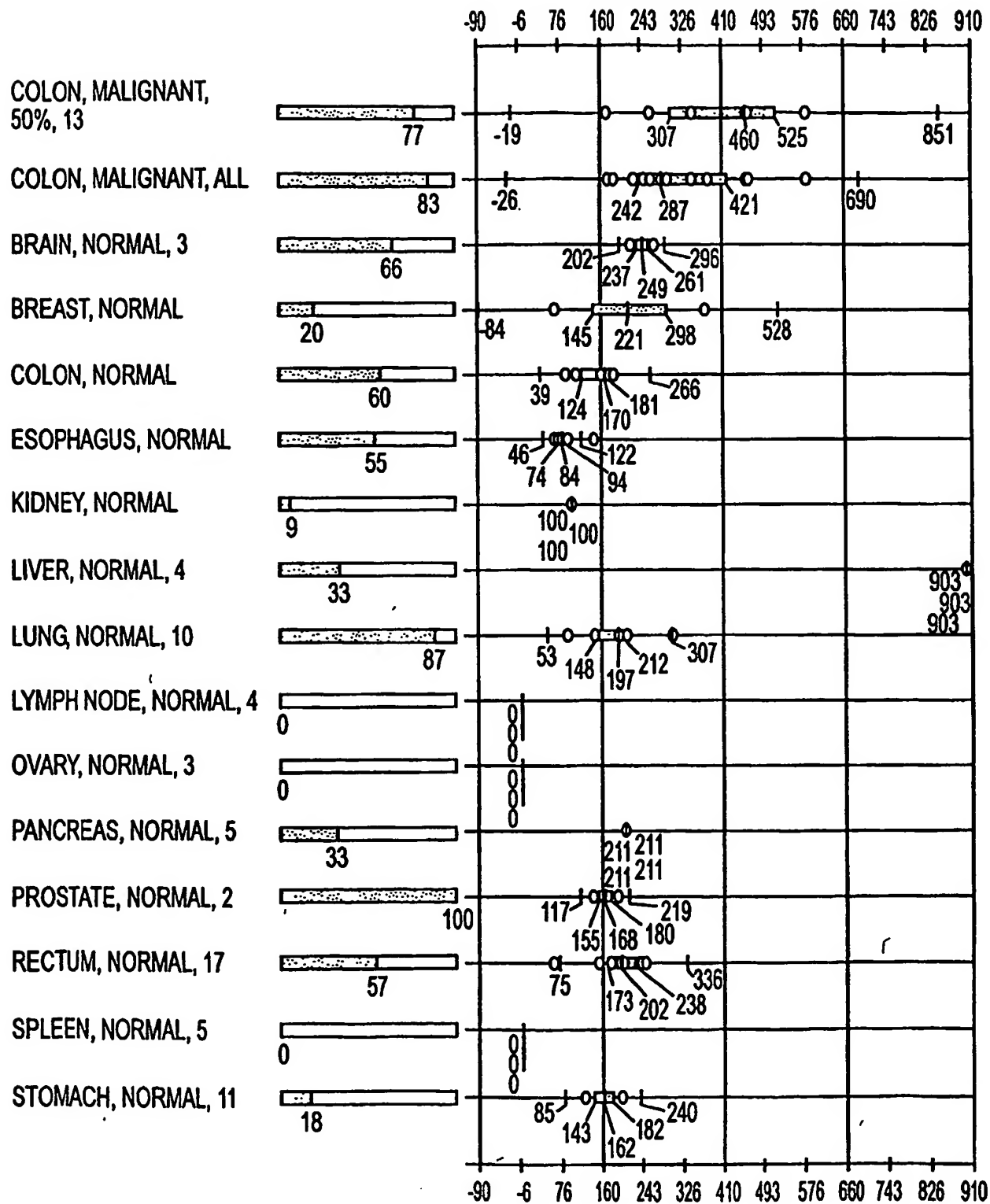
AFFY NAME : 225762\_AT

CLUSTER NAME : ESTs, WEAKLY SIMILAR TO A2AA RAT ALPHA-2A  
ADRENERGIC RECEPTOR [R.NORVEGICUS]**FIG 14**

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AFFY NAME : 229139\_AT

CLUSTER NAME : ESTs

**FIG 15**

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AFFY NAME : 232504\_AT

CLUSTER NAME : HOMO SAPIENS mRNA FULL LENGTH INSERT cDNA

CLONE EUROIMAGE 2005635

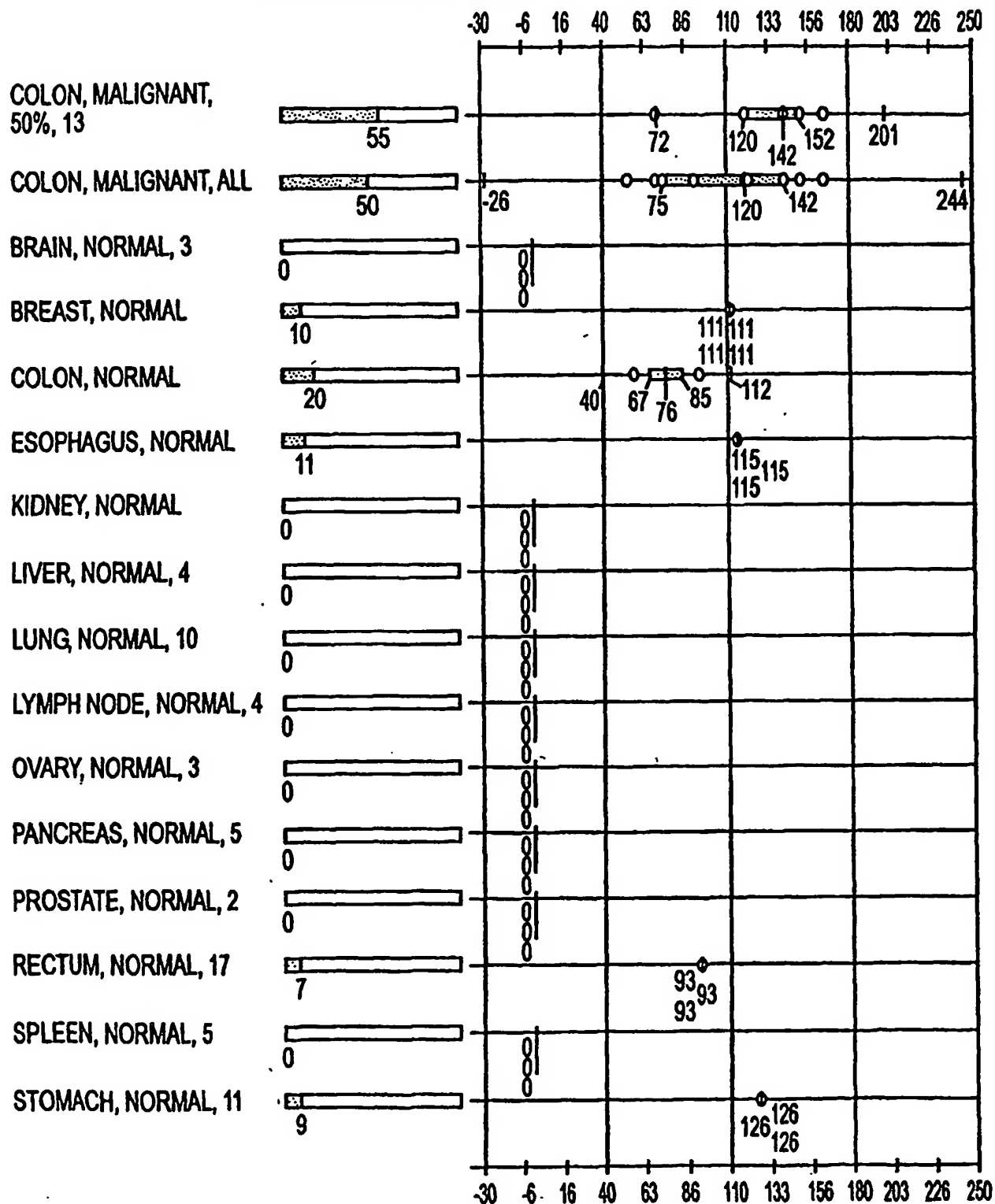


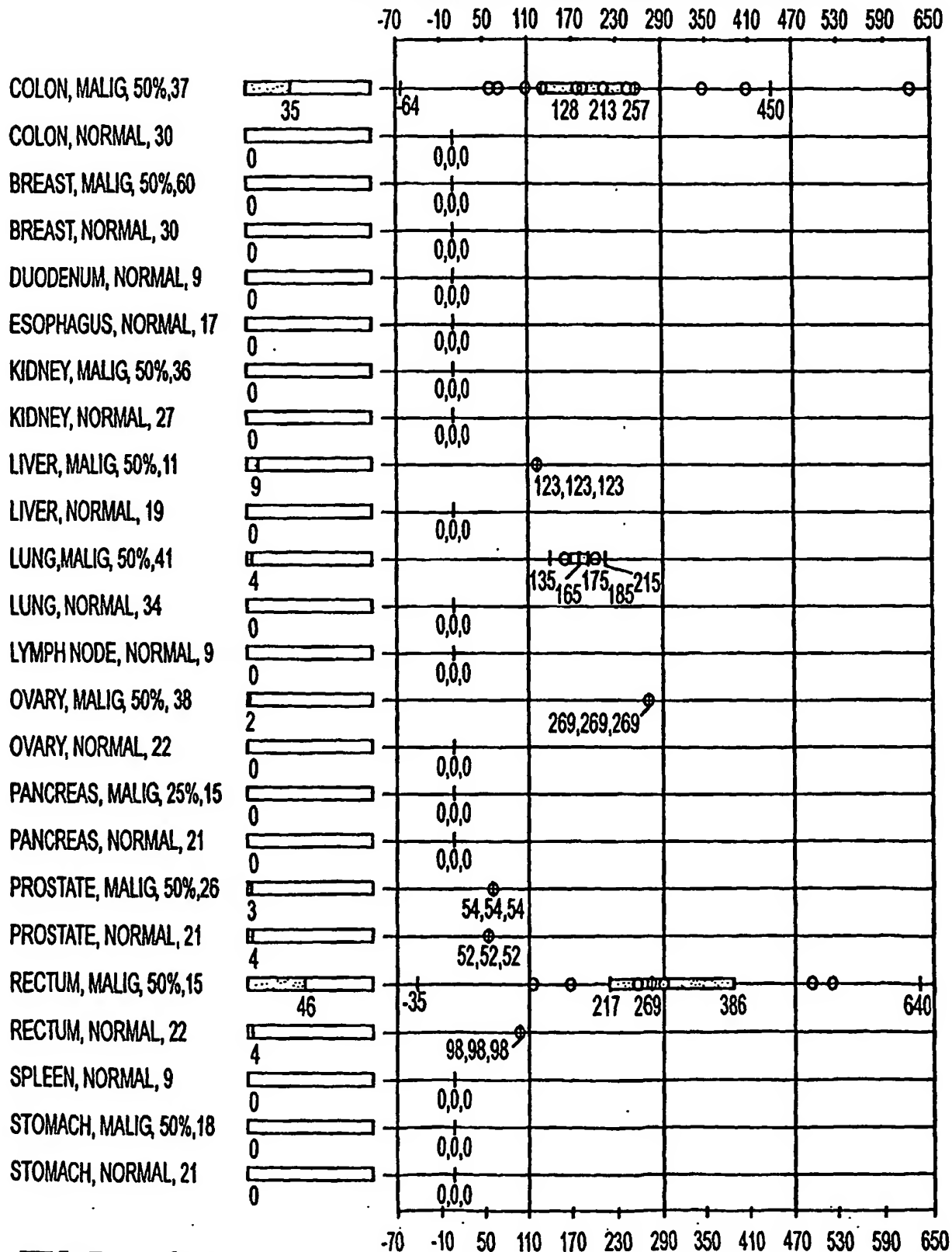
FIG. 16.



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AFFY NAME : 207457 S AT

CLUSTER NAME : CHROMOSOME 6 OPEN READING FRAME 23

**FIG. 17**

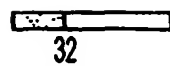
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AFFY NAME : 230615 AT

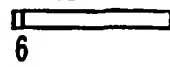
CLUSTER NAME : HOMO SAPIENS CDNA FL J32334 FIS, CLONE PROST2005426

-280 -94 91 277 463 549 835 1020 1206 1392 1578 1764 1950

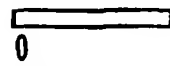
COLON, MALIG, 50%,37



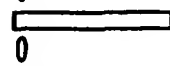
COLON, NORMAL, 30



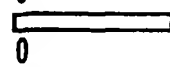
BREAST, MALIG, 50%,60



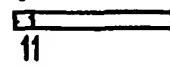
BREAST, NORMAL, 30



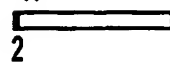
DUODENUM, NORMAL, 9



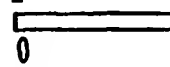
ESOPHAGUS, NORMAL, 17



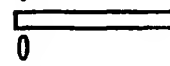
KIDNEY, MALIG, 50%,36



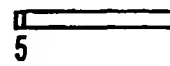
KIDNEY, NORMAL, 27



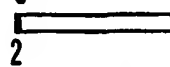
LIVER, MALIG, 50%,11



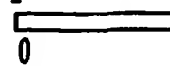
LIVER, NORMAL, 19



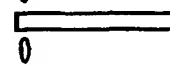
LUNG, MALIG, 50%,41



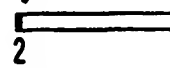
LUNG, NORMAL, 34



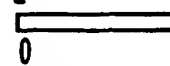
LYMPH NODE, NORMAL, 9



OVARY, MALIG, 50%, 38



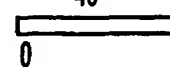
OVARY, NORMAL, 22



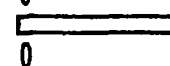
PANCREAS, MALIG, 25%,15



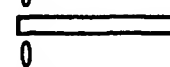
PANCREAS, NORMAL, 21



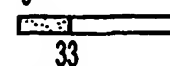
PROSTATE, MALIG, 50%,26



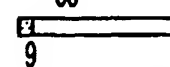
PROSTATE, NORMAL, 21



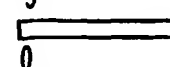
RECTUM, MALIG, 50%,15



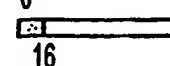
RECTUM, NORMAL, 22



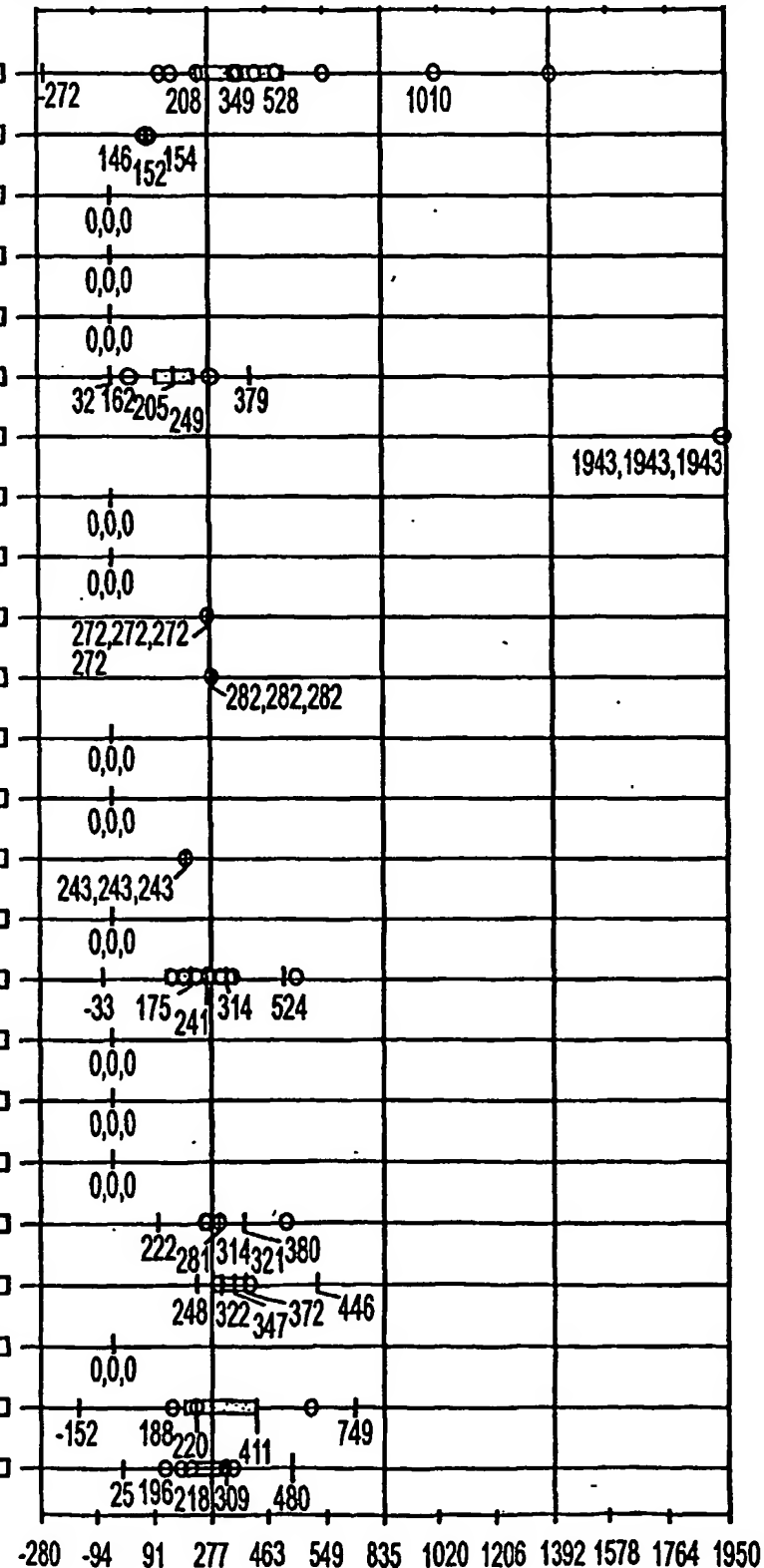
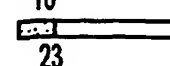
SPLEEN, NORMAL, 9



STOMACH, MALIG, 50%,18



STOMACH, NORMAL, 21

**FIG. 18**

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